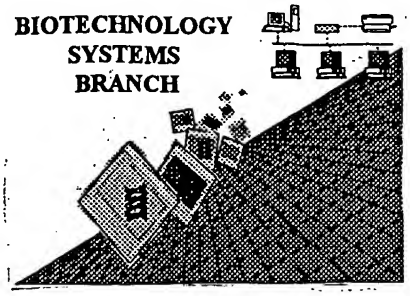
5040 #
CW 3



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/018,445
Source: PCT/IB
Date Processed by STIC: 1/14/2002

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PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
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Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom, including:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , **EFS Submission User Manual** - ePAVE)
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PCT10

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/018,445

DATE: 01/14/2002
TIME: 07:44:23

Input Set : A:\Sequence
Output Set: N:\CRF3\01142002\J018445.raw

jpr 1/14-5

<110> APPLICANT: C. Frank Bennett
Brett P. Monia
Lex M. Cowsert
ISIS PHARMACEUTICALS, INC.

<120> TITLE OF INVENTION: ANTISENSE MODULATION OF INTEGRIN BETA 3 EXPRESSION

<130> FILE REFERENCE: RTSP-0047

E--> **<140> CURRENT APPLICATION NUMBER: US/10/018,445**

E--> **<141> CURRENT FILING DATE: 2001-12-13**

<150> PRIOR APPLICATION NUMBER: US 09/344,520

<151> PRIOR FILING DATE: 1999-06-25

Does Not Comply
Corrected Diskette Needed

<160> NUMBER OF SEQ ID NOS: 47

<210> SEQ ID NO: 1
<211> LENGTH: 3170
<212> TYPE: DNA
<213> ORGANISM: Homo sapiens
<220> FEATURE:
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<222> LOCATION: (21)..(2387)
<400> SEQUENCE: 1

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   tgg gtg act gtg ctg gcg ctg ggg gcg ctg gcg ggc gtt ggc gta gga       98
   Trp Val Thr Val Leu Ala Leu Gly Ala Leu Ala Gly Val Gly Val Gly
                    15                  20                  25
   ggg ccc aac atc tgt acc acg cga ggt gtg agc tcc tgc cag cag tgc      146
   Gly Pro Asn Ile Cys Thr Thr Arg Gly Val Ser Ser Cys Gln Gln Cys
                30                  35                  40
   ctg gct gtg agc ccc atg tgt gcc tgg tgc tct gat gag gcc ctg cct      194
   Leu Ala Val Ser Pro Met Cys Ala Trp Cys Ser Asp Glu Ala Leu Pro
            45                  50                  55
   ctg ggc tca cct cgc tgt gac ctg aag gag aat ctg ctg aag gat aac      242
   Leu Gly Ser Pro Arg Cys Asp Leu Lys Glu Asn Leu Leu Lys Asp Asn
         60                  65                  70
   tgt gcc cca gaa tcc atc gag ttc cca gtg agt gag gcc cga gta cta      290
   Cys Ala Pro Glu Ser Ile Glu Phe Pro Val Ser Glu Ala Arg Val Leu
    75                  80                  85                  90
   gag gac agg ccc ctc agc gac aag ggc tct gga gac agc tcc cag gtc      338
   Glu Asp Arg Pro Leu Ser Asp Lys Gly Ser Gly Asp Ser Ser Gln Val
                   95                 100                 105
   act caa gtc agt ccc cag agg att gca ctc cgg ctc cgg cca gat gat      386
   Thr Gln Val Ser Pro Gln Arg Ile Ala Leu Arg Leu Arg Pro Asp Asp
W--> 58          110 110             115 115             120 120
   tcg aag aat ttc tcc atc caa gtg cgg cag gtg gag gat tac cct gtg      434
   Ser Lys Asn Phe Ser Ile Gln Val Arg Gln Val Glu Asp Tyr Pro Val
           125                 130                 135
   gac atc tac tac ttg atg gac ctg tct tac tcc atg aag gat gat ctg      482
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fix amino acid numbering

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/018,445

DATE: 01/14/2002
TIME: 07:44:23

Input Set : A:\Sequence
Output Set: N:\CRF3\01142002\J018445.raw

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65    Asp Ile Tyr Tyr Leu Met Asp Leu Ser Tyr Ser Met Lys Asp Asp Leu
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68    tgg agc atc cag aac ctg ggt acc aag ctg gcc acc cag atg cga aag     530
69    Trp Ser Ile Gln Asn Leu Gly Thr Lys Leu Ala Thr Gln Met Arg Lys
70    155                 160                 165                 170
72    ctc acc agt aac ctg cgg att ggc ttc ggg gca ttt gtg gac aag cct     578
73    Leu Thr Ser Asn Leu Arg Ile Gly Phe Gly Ala Phe Val Asp Lys Pro
74                    175                 180                 185
76    gtg tca cca tac atg tat atc tcc cca cca gag gcc ctc gaa aac ccc     626
77    Val Ser Pro Tyr Met Tyr Ile Ser Pro Pro Glu Ala Leu Glu Asn Pro
78                190                 195                 200
80    tgc tat gat atg aag acc acc tgc ttg ccc atg ttt ggc tac aaa cac     674
81    Cys Tyr Asp Met Lys Thr Thr Cys Leu Pro Met Phe Gly Tyr Lys His
82            205                 210                 215
84    gtg ctg acg cta act gac cag gtg acc cgc ttc aat gag gaa gtg aag     722
85    Val Leu Thr Leu Thr Asp Gln Val Thr Arg Phe Asn Glu Glu Val Lys
86        220                 225                 230
88    aag cag agt gtg tca cgg aac cga gat gcc cca gag ggt ggc ttt gat     770
89    Lys Gln Ser Val Ser Arg Asn Arg Asp Ala Pro Glu Gly Gly Phe Asp
90    235                 240                 245                 250
92    gcc atc atg cag gct aca gtc tgt gat gaa aag att ggc tgg agg aat     818
93    Ala Ile Met Gln Ala Thr Val Cys Asp Glu Lys Ile Gly Trp Arg Asn
94                    255                 260                 265
96    gat gca tcc cac ttg ctg gtg ttt acc act gat gcc aag act cat ata     866
97    Asp Ala Ser His Leu Leu Val Phe Thr Thr Asp Ala Lys Thr His Ile
98                270                 275                 280
100    gca ttg gac gga agg ctg gca ggc att gtc cag cct aat gac ggg cag     914
101    Ala Leu Asp Gly Arg Leu Ala Gly Ile Val Gln Pro Asn Asp Gly Gln
102            285                 290                 295
104    tgt cat gtt ggt agt gac aat cat tac tct gcc tcc act acc atg gat     962
105    Cys His Val Gly Ser Asp Asn His Tyr Ser Ala Ser Thr Thr Met Asp
106        300                 305                 310
108    tat ccc tct ttg ggg ctg atg act gag aag cta tcc cag aaa aac atc    1010
109    Tyr Pro Ser Leu Gly Leu Met Thr Glu Lys Leu Ser Gln Lys Asn Ile
110    315                 320                 325                 330
112    aat ttg atc ttt gca gtg act gaa aat gta gtc aat ctc tat cag aac    1058
113    Asn Leu Ile Phe Ala Val Thr Glu Asn Val Val Asn Leu Tyr Gln Asn
114                    335                 340                 345
116    tat agt gag ctc atc cca ggg acc aca gtt ggg gtt ctg tcc atg gat    1106
117    Tyr Ser Glu Leu Ile Pro Gly Thr Thr Val Gly Val Leu Ser Met Asp
118                350                 355                 360
120    tcc agc aat gtc ctc cag ctc att gtt gat gct tat ggg aaa atc cgt    1154
121    Ser Ser Asn Val Leu Gln Leu Ile Val Asp Ala Tyr Gly Lys Ile Arg
122            365                 370                 375
124    tct aaa gtc gag ctg gaa gtg cgt gac ctc cct gaa gag ttg tct cta    1202
125    Ser Lys Val Glu Leu Glu Val Arg Asp Leu Pro Glu Glu Leu Ser Leu
126        380                 385                 390
128    tcc ttc aat gcc acc tgc ctc aac aat gag gtc atc cct ggc ctc aag    1250
129    Ser Phe Asn Ala Thr Cys Leu Asn Asn Glu Val Ile Pro Gly Leu Lys
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RAW SEQUENCE LISTING DATE: 01/14/2002
PATENT APPLICATION: US/10/018,445 TIME: 07:44:23

Input Set : A:\Sequence
Output Set: N:\CRF3\01142002\J018445.raw

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132    tct tgt atg gga ctc aag att gga gac acg gtg agc ttc agc att gag    1298
133    Ser Cys Met Gly Leu Lys Ile Gly Asp Thr Val Ser Phe Ser Ile Glu
134                    415                 420                 425
136    gcc aag gtg cga ggc tgt ccc cag gag aag gag aag tcc ttt acc ata    1346
137    Ala Lys Val Arg Gly Cys Pro Gln Glu Lys Glu Lys Ser Phe Thr Ile
138                430                 435                 440
140    aag ccc gtg ggc ttc aag gac agc ctg atc gtc cag gtc acc ttt gat    1394
141    Lys Pro Val Gly Phe Lys Asp Ser Leu Ile Val Gln Val Thr Phe Asp
142            445                 450                 455
144    tgt gac tgt gcc tgc cag gcc caa gct gaa cct aat agc cat cgc tgc    1442
145    Cys Asp Cys Ala Cys Gln Ala Gln Ala Glu Pro Asn Ser His Arg Cys
146        460                 465                 470
148    aac aat ggc aat ggg acc ttt gag tgt ggg gta tgc cgt tgt ggg cct    1490
149    Asn Asn Gly Asn Gly Thr Phe Glu Cys Gly Val Cys Arg Cys Gly Pro
150    475                 480                 485                 490
152    ggc tgg ctg gga tcc cag tgt gag tgc tca gag gag gac tat cgc cct    1538
153    Gly Trp Leu Gly Ser Gln Cys Glu Cys Ser Glu Glu Asp Tyr Arg Pro
154                    495                 500                 505
156    tcc cag cag gac gag tgc agc ccc cga gag ggt cag ccc gtc tgc agc    1586
157    Ser Gln Gln Asp Glu Cys Ser Pro Arg Glu Gly Gln Pro Val Cys Ser
158                510                 515                 520
160    cag cgg ggc gag tgc ctc tgt ggt caa tgt gtc tgc cac agc agt gac    1634
161    Gln Arg Gly Glu Cys Leu Cys Gly Gln Cys Val Cys His Ser Ser Asp
162            525                 530                 535
164    ttt ggc aag atc acg ggc aag tac tgc gag tgt gac gac ttc tcc tgt    1682
165    Phe Gly Lys Ile Thr Gly Lys Tyr Cys Glu Cys Asp Asp Phe Ser Cys
166        540                 545                 550
168    gtc cgc tac aag ggg gag atg tgc tca ggc cat ggc cag tgc agc tgt    1730
169    Val Arg Tyr Lys Gly Glu Met Cys Ser Gly His Gly Gln Cys Ser Cys
170    555                 560                 565                 570
172    ggg gac tgc ctg tgt gac tcc gac tgg acc ggc tac tac tgc aac tgt    1778
173    Gly Asp Cys Leu Cys Asp Ser Asp Trp Thr Gly Tyr Tyr Cys Asn Cys
174                    575                 580                 585
176    acc acg cgt act gac acc tgc atg tcc agc aat ggg ctg ctg tgc agc    1826
177    Thr Thr Arg Thr Asp Thr Cys Met Ser Ser Asn Gly Leu Leu Cys Ser
178                590                 595                 600
180    ggc cgc ggc aag tgt gaa tgt ggc agc tgt gtc tgt atc cag ccg ggc    1874
181    Gly Arg Gly Lys Cys Glu Cys Gly Ser Cys Val Cys Ile Gln Pro Gly
182            605                 610                 615
184    tcc tat ggg gac acc tgt gag aag tgc ccc acc tgc cca gat gcc tgc    1922
185    Ser Tyr Gly Asp Thr Cys Glu Lys Cys Pro Thr Cys Pro Asp Ala Cys
186        620                 625                 630
188    acc ttt aag aaa gaa tgt gtg gag tgt aag aag ttt gac cgg gag ccc    1970
189    Thr Phe Lys Lys Glu Cys Val Glu Cys Lys Lys Phe Asp Arg Glu Pro
190    635                 640                 645                 650
192    tac atg acc gaa aat acc tgc aac cgt tac tgc cgt gac gag att gag    2018
193    Tyr Met Thr Glu Asn Thr Cys Asn Arg Tyr Cys Arg Asp Glu Ile Glu
194                    655                 660                 665
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RAW SEQUENCE LISTING DATE: 01/14/2002
PATENT APPLICATION: US/10/018,445 TIME: 07:44:23

Input Set : A:\Sequence
Output Set: N:\CRF3\01142002\J018445.raw

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196    tca gtg aaa gag ctt aag gac act ggc aag gat gca gtg aat tgt acc    2066
197    Ser Val Lys Glu Leu Lys Asp Thr Gly Lys Asp Ala Val Asn Cys Thr
198                670                 675                 680
200    tat aag aat gag gat gac tgt gtc gtc aga ttc cag tac tat gaa gat    2114
201    Tyr Lys Asn Glu Asp Asp Cys Val Val Arg Phe Gln Tyr Tyr Glu Asp
202            685                 690                 695
204    tct agt gga aag tcc atc ctg tat gtg gta gaa gag cca gag tgt ccc    2162
205    Ser Ser Gly Lys Ser Ile Leu Tyr Val Val Glu Glu Pro Glu Cys Pro
206        700                 705                 710
208    aag ggc cct gac atc ctg gtg gtc ctg ctc tca gtg atg ggg gcc att    2210
209    Lys Gly Pro Asp Ile Leu Val Val Leu Leu Ser Val Met Gly Ala Ile
210    715                 720                 725                 730
212    ctg ctc att ggc ctt gcc gcc ctg ctc atc tgg aaa ctc ctc atc acc    2258
213    Leu Leu Ile Gly Leu Ala Ala Leu Leu Ile Trp Lys Leu Leu Ile Thr
214                735                 740                 745
216    atc cac gac cga aaa gaa ttc gct aaa ttt gag gaa gaa cgc gcc aga    2306
217    Ile His Asp Arg Lys Glu Phe Ala Lys Phe Glu Glu Glu Arg Ala Arg
218                750                 755                 760
220    gca aaa tgg gac aca gcc aac aac cca ctg tat aaa gag gcc acg tct    2354
221    Ala Lys Trp Asp Thr Ala Asn Asn Pro Leu Tyr Lys Glu Ala Thr Ser
222            765                 770                 775
224    acc ttc acc aat atc acg tac cgg ggc act taa tgataagcag tcatcctcag  2407
225    Thr Phe Thr Asn Ile Thr Tyr Arg Gly Thr
226        780                 785
228    atcattatca gcctgtgcca ggattgcagg agtccctgcc atcatgttta cagaggacag  2467
230    tatttgtggg gagggatttc ggggctcaga gtggggtagg ttgggagaat gtcagtatgt  2527
232    ggaagtgtgg gtctgtgtgt gtgtatgtgg gggtctgtgt gtttatgtgt gtgtgttgtg  2587
234    tgtgggagtg tgtaatttaa aattgtgatg tgtcctgata agctgagctc cttagccttt  2647
236    gtcccagaat gcctcctgca gggattcttc ctgcttagct tgagggtgac tatggagctg  2707
238    agcaggtgtt cttcattacc tcagtgagaa gccagctttc ctcatcaggc cattgtccct  2767
240    gaagagaagg gcagggctga ggcctctcat tccagaggaa gggacaccaa gccttggctc  2827
242    taccctgagt tcataaattt atggttctca ggcctgactc tcagcagcta tggtaggaac  2887
244    tgctggcttg gcagcccggg tcatctgtac ctctgcctcc tttcccctcc ctcaggccga  2947
246    aggaggagtc agggagagct gaactattag agctgcctgt gccttttgcc atcccctcaa  3007
248    cccagctatg gttctctcgc aagggaagtc cttgcaagct aattctttga cctgttggga  3067
250    gtgaggatgt ctgggccact caggggtcat tcatggcctg ggggatgtac cagcatctcc  3127
252    cagttcataa tcacaaccct tcagatttgc cttattggca gcg                    3170
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256 <211> LENGTH: 23
257 <212> TYPE: DNA
258 <213> ORGANISM: Artificial Sequence
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266 <210> SEQ ID NO: 3
267 <211> LENGTH: 21
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269 <213> ORGANISM: Artificial Sequence
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insert this mandatory numeric identifier wherever <2217, <2227, or <2237 is shown (global error)

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/018,445

DATE: 01/14/2002
TIME: 07:44:23

Input Set : A:\Sequence
Output Set: N:\CRF3\01142002\J018445.raw

same error

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271 <223> OTHER INFORMATION: PCR Primer
273 <400> SEQUENCE: 3
274 ccgtcattag gctggacaat g 21
277 <210> SEQ ID NO: 4
278 <211> LENGTH: 25
279 <212> TYPE: DNA
280 <213> ORGANISM: Artificial Sequence
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288 <210> SEQ ID NO: 5
289 <211> LENGTH: 19
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291 <213> ORGANISM: Artificial Sequence
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299 <210> SEQ ID NO: 6
300 <211> LENGTH: 20
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302 <213> ORGANISM: Artificial Sequence
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313 <213> ORGANISM: Artificial Sequence
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323 <212> TYPE: DNA
324 <213> ORGANISM: Artificial Sequence
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334 <212> TYPE: DNA
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The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

VERIFICATION SUMMARY DATE: 01/14/2002
PATENT APPLICATION: US/10/018,445 TIME: 07:44:24

Input Set : A:\Sequence
Output Set: N:\CRF3\01142002\J018445.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application No
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:58 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:260 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:271 M:258 W: Mandatory Feature missing, <220> FEATURE:
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VERIFICATION SUMMARY
PATENT APPLICATION: **US/10/018,445**

DATE: 01/14/2002
TIME: 07:44:24

Input Set : **A:\Sequence**
Output Set: **N:\CRF3\01142002\J018445.raw**

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